

#6

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/887,038

DATE: 10/04/2001
TIME: 08:07:45

Input Set : A:\SEQUENCE LISTING.txt
Output Set: N:\CRF3\10042001\I887038.raw

5 <110> APPLICANT: Kaplan, Aaron
7 Lieman-Hurwitz, Judy
9 Schatz, Daniella
11 Mittler, Ron
13 Ronen-Tarazi, Michal
15 Bonfil, David J.
19 <120> TITLE OF INVENTION: ENHANCING INORGANIC CARBON FIXATION BY PHOTOSYNTHETIC
ORGANISMS

23 <130> FILE REFERENCE: 01/22171
C--> 27 <140> CURRENT APPLICATION NUMBER: US/09/887,038

C--> 27 <141> CURRENT FILING DATE: 2001-09-17

27 <160> NUMBER OF SEQ ID NOS: 9

31 <170> SOFTWARE: PatentIn version 3.1

35 <210> SEQ ID NO: 1

37 <211> LENGTH: 4957

39 <212> TYPE: DNA

41 <213> ORGANISM: Synechococcus sp.

45 <400> SEQUENCE: 1

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277 <212> TYPE: PRT
279 <213> ORGANISM: Synechococcus sp.
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290 20 25 30
293 Trp Arg Ala Ser Ser Gln Leu Leu Val Trp Ser Glu Ala Leu Gly Gly
294 35 40 45
297 Phe Leu Ala Val Val Tyr Gly Ser Ala Pro Phe Val Pro Ser Ser
298 50 55 60
301 Ala Leu Gly Leu Gly Leu Ala Ala Ile Ala Ala Tyr Trp Ala Leu Leu
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305 Ser Leu Thr Asp Ile Asp Leu Arg Gln Ala Thr Pro Ile His Trp Leu

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310	100	105	110	
313	Val Arg Ala Ala Ala Leu Val Gly Leu Ala Lys Leu Thr Leu Tyr Leu			
314	115	120	125	
317	Leu Val Phe Ala Leu Ala Ala Arg Val Leu Arg Asn Pro Arg Leu Arg			
318	130	135	140	
321	Ser Leu Leu Phe Ser Val Val Val Ile Thr Ser Leu Phe Val Ser Val			
322	145	150	155	160
325	Tyr Gly Leu Asn Gln Trp Ile Tyr Gly Val Glu Glu Leu Ala Thr Trp			
326	165	170	175	
329	Val Asp Arg Asn Ser Val Ala Asp Phe Thr Ser Arg Val Tyr Ser Tyr			
330	180	185	190	
333	Leu Gly Asn Pro Asn Leu Leu Ala Ala Tyr Leu Val Pro Thr Thr Ala			
334	195	200	205	
337	Phe Ser Ala Ala Ala Ile Gly Val Trp Arg Gly Trp Leu Pro Lys Leu			
338	210	215	220	
341	Leu Ala Ile Ala Ala Thr Gly Ala Ser Ser Leu Cys Leu Ile Leu Thr			
342	225	230	235	240
345	Tyr Ser Arg Gly Gly Trp Leu Gly Phe Val Ala Met Ile Phe Val Trp			
346	245	250	255	
349	Ala Leu Leu Gly Leu Tyr Trp Phe Gln Pro Arg Leu Pro Ala Pro Trp			
350	260	265	270	
353	Arg Arg Trp Leu Phe Pro Val Val Leu Gly Gly Leu Val Ala Val Leu			
354	275	280	285	
357	Leu Val Ala Val Leu Gly Leu Glu Pro Leu Arg Val Arg Val Leu Ser			
358	290	295	300	
361	Ile Phe Val Gly Arg Glu Asp Ser Ser Asn Asn Phe Arg Ile Asn Val			
362	305	310	315	320
365	Trp Leu Ala Val Leu Gln Met Ile Gln Asp Arg Pro Trp Leu Gly Ile			
366	325	330	335	
369	Gly Pro Gly Asn Thr Ala Phe Asn Leu Val Tyr Pro Leu Tyr Gln Gln			
370	340	345	350	
373	Ala Arg Phe Thr Ala Leu Ser Ala Tyr Ser Val Pro Leu Glu Val Ala			
374	355	360	365	
377	Val Glu Gly Gly Leu Leu Gly Leu Thr Ala Phe Ala Trp Leu Leu Leu			
378	370	375	380	
381	Val Thr Ala Val Thr Ala Val Arg Gln Val Ser Arg Leu Arg Arg Asp			
382	385	390	395	400
385	Arg Asn Pro Gln Ala Phe Trp Leu Met Ala Ser Leu Ala Gly Leu Ala			
386	405	410	415	
389	Gly Met Leu Gly His Gly Leu Phe Asp Thr Val Leu Tyr Arg Pro Glu			
390	420	425	430	
393	Ala Ser Thr Leu Trp Trp Leu Cys Ile Gly Ala Ile Ala Ser Phe Trp			
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397	Gln Pro Gln Pro Ser Lys Gln Leu Pro Pro Glu Ala Glu His Ser Asp			
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405 <210> SEQ ID NO: 4
407 <211> LENGTH: 1425
409 <212> TYPE: DNA
411 <213> ORGANISM: Synechococcus sp.
415 <400> SEQUENCE: 4
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486 35 40 45
489 Ala Leu Gly Thr Ala Leu Val Ala Ile Ile Phe Ile Ala Ala Pro Phe
490 50 55 60
493 Thr Ser Thr Thr Met Leu Gly Ile Phe Met Leu Leu Cys Gly Ala Phe
494 65 70 75 80
497 Trp Ala Leu Leu Thr Phe Ala Asp Gln Pro Gly Lys Gly Leu Thr Pro
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502 100 105 110
505 Gly Phe Ser Pro Val Lys Met Ala Ala Ser Gly Leu Ala Lys Leu

VERIFICATION SUMMARY
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L:27 M:270 C: Current Application Number differs, Replaced Current Application No
L:27 M:271 C: Current Filing Date differs, Replaced Current Filing Date